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STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 101578, S21

Source: TECUBP

Date Processed by STIC: S-17-06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)**
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/10/06

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Raw Sequence Listing Error Summary

ERROR DETECTED**SUGGESTED CORRECTION****SERIAL NUMBER:**

101578,521

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
(OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per I.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
Response Per I.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>

Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. I.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

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IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006

TIME: 10:12:56

Input Set : A:\50413.015001.txt
 Output Set: N:\CRF4\05172006\J578521.raw

3 <110> APPLICANT: Chun, Jong-Yoon
 5 <120> TITLE OF INVENTION: Method for Amplifying Unknown DNA Sequence Adjacent to Known
 6 Sequence
 8 <130> FILE REFERENCE: 50413/015001
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/578,521
 C--> 10 <141> CURRENT FILING DATE: 2006-05-08
 10 <150> PRIOR APPLICATION NUMBER: PCT/KR2003/002407
 11 <151> PRIOR FILING DATE: 2003-11-10
 13 <160> NUMBER OF SEQ ID NOS: 28
 15 <170> SOFTWARE: PatentIn version 3.3
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 33
 19 <212> TYPE: DNA
 20 <213> ORGANISM: Artificial Sequence *Pls explain source of genetic material.* Does Not Use Corrected Diskette Needed (Pg. 1-5) 2
 22 <220> FEATURE:
 23 <223> OTHER INFORMATION: DW-ACP1-A *INVALID RESPONSE*
 26 <220> FEATURE:
 27 <221> NAME/KEY: misc_feature
 28 <222> LOCATION: (22)..(28)
 29 <223> OTHER INFORMATION: n denotes deoxyinosine
 31 <400> SEQUENCE: 1
 W--> 32 tcacagaagt atgccaagcg annnnnnnnag gtc
 35 <210> SEQ ID NO: 2
 36 <211> LENGTH: 33
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Artificial Sequence *Pls explain source of genetic material* 33
 40 <220> FEATURE:
 41 <223> OTHER INFORMATION: DW-ACP1-C *INVALID RESPONSE*
 44 <220> FEATURE:
 45 <221> NAME/KEY: misc_feature
 46 <222> LOCATION: (22)..(28)
 47 <223> OTHER INFORMATION: n denotes deoxyinosine
 49 <400> SEQUENCE: 2
 W--> 50 tcacagaagt atgccaagcg annnnnnnncg gtc
 53 <210> SEQ ID NO: 3
 54 <211> LENGTH: 33
 55 <212> TYPE: DNA
 56 <213> ORGANISM: Artificial Sequence
 58 <220> FEATURE:
 59 <223> OTHER INFORMATION: DW-ACP1-T *Same error* #11
 62 <220> FEATURE:
 63 <221> NAME/KEY: misc_feature
 64 <222> LOCATION: (22)..(28)

~ See item #11
 on error summary sheet 5/17/2006

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:56

Input Set : A:\50413.015001.txt
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```

65 <223> OTHER INFORMATION: n denotes deoxyinosine
67 <400> SEQUENCE: 3
W--> 68 tcacagaagt atgccaagcg annnnnnnntg gtc
71 <210> SEQ ID NO: 4
72 <211> LENGTH: 33
73 <212> TYPE: DNA
74 <213> ORGANISM: Artificial Sequence
76 <220> FEATURE:
77 <223> OTHER INFORMATION: DW-ACP1-G
80 <220> FEATURE:
81 <221> NAME/KEY: misc_feature
82 <222> LOCATION: (22)..(28)
83 <223> OTHER INFORMATION: n denotes deoxyinosine
85 <400> SEQUENCE: 4
W--> 86 tcacagaagt atgccaagcg annnnnnnngg gtc
89 <210> SEQ ID NO: 5
90 <211> LENGTH: 33
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: DW-ACP-2
98 <220> FEATURE:
99 <221> NAME/KEY: misc_feature
100 <222> LOCATION: (26)..(29)
101 <223> OTHER INFORMATION: n denotes deoxyinosine
103 <400> SEQUENCE: 5
W--> 104 tcacagaagt atgccaagcg aggggnnnng gtc
107 <210> SEQ ID NO: 6
108 <211> LENGTH: 33
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: DW-ACP2-NA
116 <220> FEATURE:
117 <221> NAME/KEY: misc_feature
118 <222> LOCATION: (26)..(28)
119 <223> OTHER INFORMATION: n denotes deoxyinosine
121 <400> SEQUENCE: 6
W--> 122 tcacagaagt atgccaagcg aggggnnnnag gtc
125 <210> SEQ ID NO: 7
126 <211> LENGTH: 33
127 <212> TYPE: DNA
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: DW-ACP2-NC
134 <220> FEATURE:
135 <221> NAME/KEY: misc_feature
136 <222> LOCATION: (26)..(28)
137 <223> OTHER INFORMATION: n denotes deoxyinosine

```

33 Same errors

See item
11 on
error
Summary
Sheet.

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:56

Input Set : A:\50413.015001.txt
Output Set: N:\CRF4\05172006\J578521.raw

33
SAME errors

139 <400> SEQUENCE: 7
W--> 140 tcacagaagt atgccaagcg aggggnncg gtc
143 <210> SEQ ID NO: 8
144 <211> LENGTH: 33
145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: DW-ACP2-NT
152 <220> FEATURE:
153 <221> NAME/KEY: misc_feature
154 <222> LOCATION: (26)..(28)
155 <223> OTHER INFORMATION: n denotes deoxyinosine
157 <400> SEQUENCE: 8
W--> 158 tcacagaagt atgccaagcg aggggnntg gtc 33
161 <210> SEQ ID NO: 9
162 <211> LENGTH: 33
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: DW-ACP2-NG
170 <220> FEATURE:
171 <221> NAME/KEY: misc_feature
172 <222> LOCATION: (26)..(28)
173 <223> OTHER INFORMATION: n denotes deoxyinosine
175 <400> SEQUENCE: 9
W--> 176 tcacagaagt atgccaagcg aggggnnnng gtc 33
179 <210> SEQ ID NO: 10
180 <211> LENGTH: 33
181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: DW-ACP3-N1
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (22)..(25)
191 <223> OTHER INFORMATION: n denotes deoxyinosine
193 <400> SEQUENCE: 10
W--> 194 tcacagaagt atgccaagcg annnnggggg gtc 33
197 <210> SEQ ID NO: 11
198 <211> LENGTH: 33
199 <212> TYPE: DNA
200 <213> ORGANISM: Artificial Sequence
202 <220> FEATURE:
203 <223> OTHER INFORMATION: DW-ACP3-N2
206 <220> FEATURE:
207 <221> NAME/KEY: misc_feature
208 <222> LOCATION: (23)..(26)
209 <223> OTHER INFORMATION: n denotes deoxyinosine
211 <400> SEQUENCE: 11

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:56

Input Set : A:\50413.015001.txt
Output Set: N:\CRF4\05172006\J578521.raw

```

W--> 212 tcacagaagt atgccaagcg agnnnnnggg gtc
215 <210> SEQ ID NO: 12
216 <211> LENGTH: 33
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: DW-ACP3-N3
224 <220> FEATURE:
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: (24)..(27)
227 <223> OTHER INFORMATION: n denotes deoxyinosine
229 <400> SEQUENCE: 12

W--> 230 tcacagaagt atgccaagcg aggnnnnggg gtc
233 <210> SEQ ID NO: 13
234 <211> LENGTH: 20
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Nested DW-P3-N
241 <400> SEQUENCE: 13
242 ccaagcggagg gggggggggc
245 <210> SEQ ID NO: 14
246 <211> LENGTH: 29
247 <212> TYPE: DNA
248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: DW-P1-A
254 <220> FEATURE:
255 <221> NAME/KEY: misc_feature
256 <222> LOCATION: (22)..(24)
257 <223> OTHER INFORMATION: n is any base
259 <400> SEQUENCE: 14

W--> 260 tcacagaagt atgccaagcg annnaggc
263 <210> SEQ ID NO: 15
264 <211> LENGTH: 29
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <223> OTHER INFORMATION: DW-P1-C
272 <220> FEATURE:
273 <221> NAME/KEY: misc_feature
274 <222> LOCATION: (22)..(24)
275 <223> OTHER INFORMATION: n is any base
277 <400> SEQUENCE: 15

W--> 278 tcacagaagt atgccaagcg annncggc
281 <210> SEQ ID NO: 16
282 <211> LENGTH: 29
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence

```

C/Same

ERRORS

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:56

Input Set : A:\50413.015001.txt
Output Set: N:\CRF4\05172006\J578521.raw

286 <220> FEATURE:
287 <223> OTHER INFORMATION: DW-P1-T
290 <220> FEATURE:
291 <221> NAME/KEY: misc_feature
292 <222> LOCATION: (22)..(24)
293 <223> OTHER INFORMATION: n is any base
295 <400> SEQUENCE: 16

↙ Same
errors

W--> 296 tcacagaagt atgccaagcg annntggtc
299 <210> SEQ ID NO: 17
300 <211> LENGTH: 29
301 <212> TYPE: DNA
302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
305 <223> OTHER INFORMATION: DW-P1-G
308 <220> FEATURE:
309 <221> NAME/KEY: misc_feature
310 <222> LOCATION: (22)..(24)
311 <223> OTHER INFORMATION: n is any base
313 <400> SEQUENCE: 17

29

See item
29 # 11 on
error
summary
sheet.

W--> 314 tcacagaagt atgccaagcg annngggc
317 <210> SEQ ID NO: 18
318 <211> LENGTH: 29
319 <212> TYPE: DNA
320 <213> ORGANISM: Artificial Sequence
322 <220> FEATURE:
323 <223> OTHER INFORMATION: DW-P1
326 <220> FEATURE:
327 <221> NAME/KEY: misc_feature
328 <222> LOCATION: (22)..(25)
329 <223> OTHER INFORMATION: n is any base
331 <400> SEQUENCE: 18

29

W--> 332 tcacagaagt atgccaagecg annnnnggc
335 <210> SEQ ID NO: 19
336 <211> LENGTH: 21
337 <212> TYPE: DNA
338 <213> ORGANISM: Artificial Sequence
340 <220> FEATURE:
341 <223> OTHER INFORMATION: JYC3
343 <400> SEQUENCE: 19
344 tcacagaagt atgccaagcg a
347 <210> SEQ ID NO: 20
348 <211> LENGTH: 20
349 <212> TYPE: DNA
350 <213> ORGANISM: Artificial Sequence
352 <220> FEATURE:
353 <223> OTHER INFORMATION: mTNFa-C1
355 <400> SEQUENCE: 20
356 caccttgcgc tgcccat tag
359 <210> SEQ ID NO: 21

21

20

The type of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:57

Input Set : A:\50413.015001.txt
Output Set: N:\CRF4\05172006\J578521.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 22,23,24,25,26,27,28
Seq#:2; N Pos. 22,23,24,25,26,27,28
Seq#:3; N Pos. 22,23,24,25,26,27,28
Seq#:4; N Pos. 22,23,24,25,26,27,28
Seq#:5; N Pos. 26,27,28
Seq#:6; N Pos. 26,27,28
Seq#:7; N Pos. 26,27,28
Seq#:8; N Pos. 26,27,28
Seq#:9; N Pos. 26,27,28
Seq#:10; N Pos. 22,23,24,25
Seq#:11; N Pos. 23,24,25,26
Seq#:12; N Pos. 24,25,26,27
Seq#:14; N Pos. 22,23,24
Seq#:15; N Pos. 22,23,24
Seq#:16; N Pos. 22,23,24
Seq#:17; N Pos. 22,23,24
Seq#:18; N Pos. 22,23,24,25
Seq#:28; N Pos. 1,2,3,4,5,6

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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/578,521

DATE: 05/17/2006
TIME: 10:12:57

Input Set : A:\50413.015001.txt
Output Set: N:\CRF4\05172006\J578521.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:32 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:68 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:158 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:260 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0